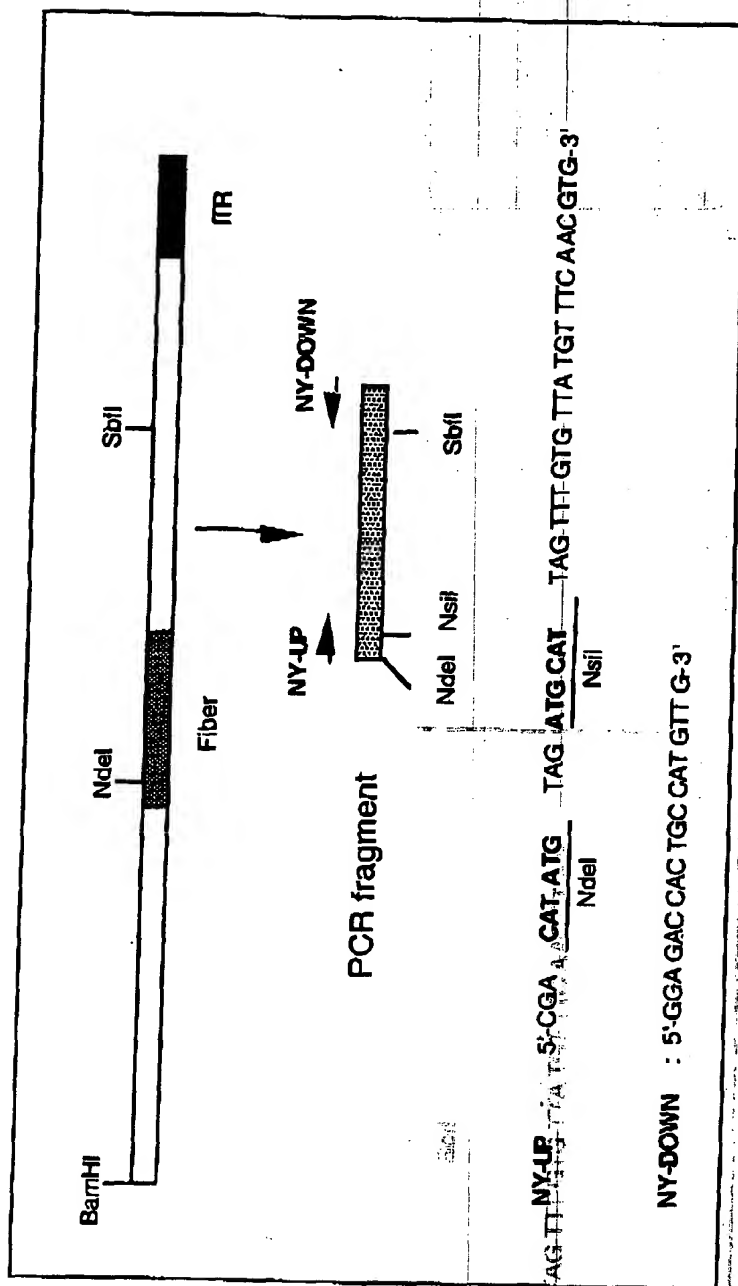


Figure 1

Figure 2



NO. 0432

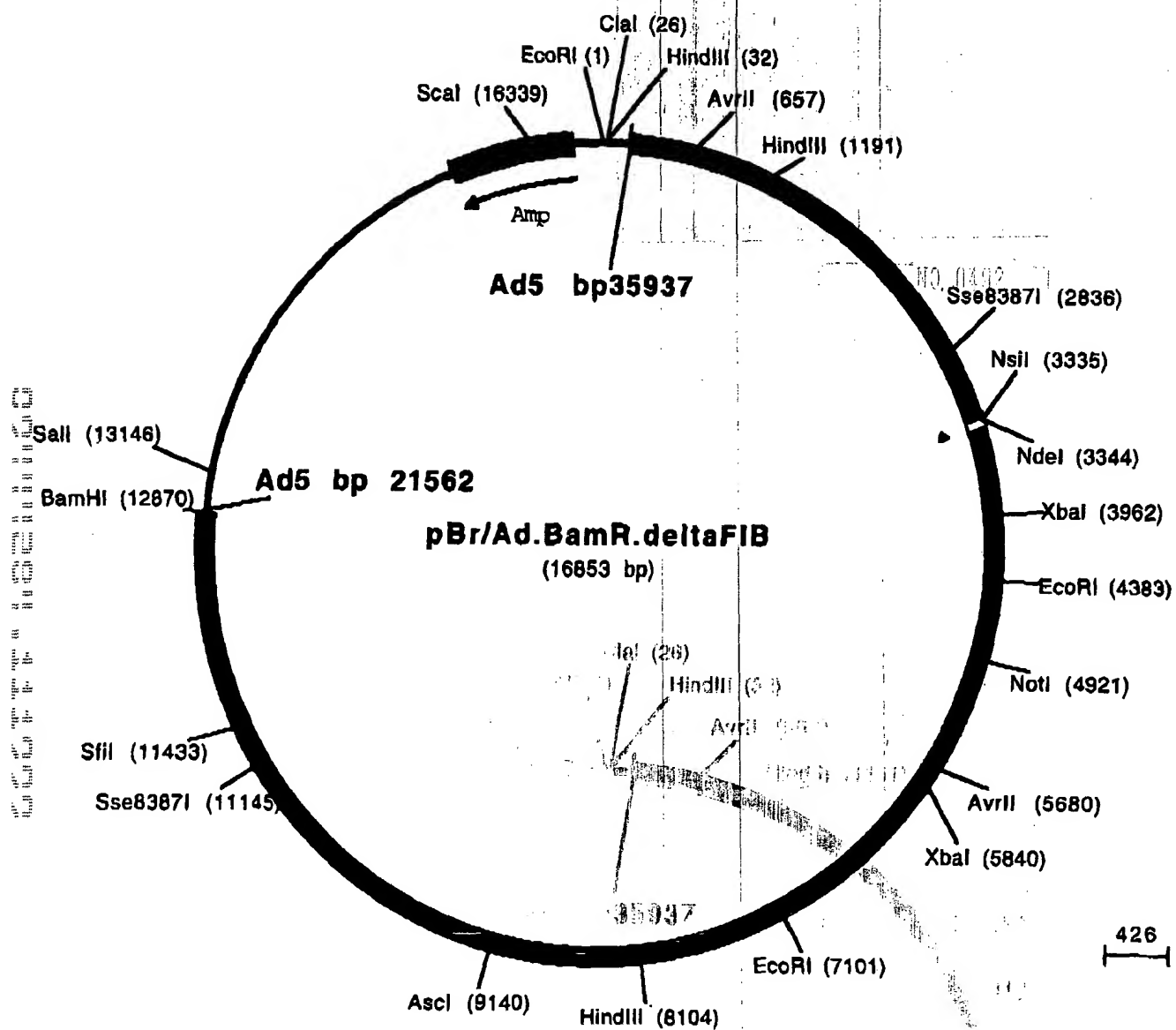


Figure 3

Figure 4A: Sequence of Ad5 fiber

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GGTACTCTCTTTGCGCCTATCCGAACCTCTAGTTACCTCCAATGGCATGCTTGCGCTCAAAATGGGC
AACGGCCTCTCTCTGGACGAGGCCGGCAACCTTACCTCCCCAAATGTAACCACTGTGAGCCCACCTC
TCAAAAAAACCAAGTCAAACATAAACCTGGAAATATCTGCACCCCTCACAGTTACCTCAGAAGCCCT
AACTGTGGCTGCCGCCGCACCTCTAATGGTCGCGGGCAACACACTCACCATGCAATCACAGGCCCCG
CTAACCGTGCACGACTCCAACTTAGCATTGCCACCCAAGGACCCCTCACAGTGTGAGAAGGAAAGC
TAGCCCTGCAAACATCAGGCCCCCTCACCACCACCGATAGCAGTACCCTTACTATCACTGCCTCACC
CCCTCTAACTACTGCCACTGGTAGCTTGGGCATTGACTTGAAAGAGCCCATTTATACACAAAATGGA
AACTAGGACTAAAGTACGGGGCTCCTTTGCATGTAACAGACGACCTAAACACTTTGACCGTAGCAA
CTGGTCCAGGTGTGACTATTAATAATACCTTCTTGCAAACATAAGTTACTGGAGCCTTGGGTTTTGA
TTCACAAGGCAATATGCAACTTAATGTAGCAGGAGGACTAAGGATTGATTCTCAAACAGACGCCTT
ATACTTGATGTTAGTTATCCGTTTGATGCTCAAACCAACTAAATCTAAGACTAGGACAGGGCCCTC
TTTTTATAAACTCAGCCCACTTGGATATTAACATAACAAAGGCCTTTACTTGTTTACAGCTTC
AAACAATTCCAAAAGCTTGAGGTTAACCTAAGCACTGCCAAGGGGTGATGTTTGACGCTACAGCC
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CCTTAGTTTTTGACAGCACAGGTGCCATTACAGTAGGAAACAAAATAATGATAAGCTAACTTTGTGG
ACCACACCAGCTCCATCTCCTAACTGTAGACTAAATGCAGAGAAAGATGCTAAACTCACTTTTGGTCT
TAACAAAATGTGGCAGTCAAATACTTGCTACAGTTTCAGTTTTGGCTGTTAAAGGCAGTTTGGCTCC
AATATCTGGAACAGTTCAAAGTGCTCATCTTATTATAAGATTTGACGAAAATGGAGTGCTACTAAC
AATTCCTTCCTGGACCCAGAATATTGGAACCTTAGAAATGGAGATCTTACTGAAGGCACAGCCTATA
CAAACGCTGTTGGATTTATGCCTAACCTATCAGCTTATCCAAATCTCACGGTAAACTGCCAAAAG
TAACATTGTCAGTCAAGTTTACTTAAACGGAGACAAAACATAACCTGTAACACTAACCATTACACTA
AACGGTACACAGGAAACAGGAGACACAACCTCCAAGTGCATCTCTATGTCATTTTCATGGGACTGGT
CTGGCCACAACCTACATTAATGAAATATTTGCCACATCTCTTACACTTTTTTCATACATTGCCCAAGA
ATAA

Figure 4B: Sequence of Ad5/fib12 chimeric fiber

ATGAAGCGCGCAAGACCGTCTGAAGATACCTTCAACCCCGTGTATCCATATGACCCATTTGACACAT
CAGACGTACCCTTTGTTACACCCCTTTTACTTCTTCCAATGGTCTTCAAGAAAACCACCAGGTGT
ATTAGCACTTAATTACAAAGACCCCATTTGTAAGTGAAGCTTGAACCCCTTACACTCAAGCTAGGGGAC
GGAATAAACTTAATGCCCCAAGGTCAACTTACAGCTAGTAATAATATCAATGTTTTGGAGCCCCTTA
CCAACACCTCACAAGGTCTTAACTTTCTTGGAGCGCCCCCTTAGCAGTAAAGGCTAGTGCCCTCAC
ACTTAACACAAGAGCGCCCTTAACCACAACGGATGAAAGCTTAGCCTTAATAACCGCCCCCTCCCATT
ACAGTAGAGTCTTCGCGTTTGGGCTTGGCCACCATAGCCCTCTAAGCTTAGATGGAGGTGGAAACC
TAGGTTTAAATCTTTCTGCTCCCCTGGACGTAGTAACAACAATTTGCATCTCACCCTGAAACTCC
CTTAGTTGTAAATTCTAGCGGTGCCCTATCTGTTGCTACTGCAGACCCCATAGTGTTCGCAACAAC
GCTCTTACCCTACCTACGGCAGATCCGTTAATGGTGAGCTCCGATGGGTGGGAATAAGTGTCACTA
GTCCCATTACAGTAATAAACGGTTCCTTAGCCTTGCTTACAACGTCTCCCCTCAACAGCACAGGATC
CACTTTAAGTCTGTCTGTTGCCAATCCTCTGACTATTTACAAGACACATTGACTGTTTCCACTGGT
AACGGTCTTCAAGTGTGCGGGTCTCAATTAGTAACAAGAATAGGGGATGGTTTAACATTCGATAATG
GGGTCATGAAAGTAAACGTTGCCGGGGGAATGAGAACTTCTGGCGGTAGAATAATTTTAGATGTTAA
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CTCTTAATGCAGGTCAGGGGCTTACATTTAATAATGGCCAACCTTAGGGTTAAGTTGGGAGCTGGACT
TATTTTTGATTCAAACAATAACATTGCCTTAGGCAGCAGCAGCAACACTCCATACGACCCTCTGACA
CTGTGGACAACCTCTGACCCACCACCAAACTGCAGCCTCATAACAAGAGCTAGATGCAAAACTCACCC
TGTGCTTAACAAAAAACGGATCTATTGTTAATGGCATTGTAAGTTTAGTGGGTGTTAAGGGTAATCT
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AATGCATTAG

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TAACAACAATTTTCTTACCACTG
CTACTGCAATCTTACCTTACCTG
TAGCTCCGATCTTACCTTACCTG
CTTACAACCTTACCTTACCTG
ATTTCACAACCTTACCTTACCTG
TAGGATACTTACCTTACCTG
ACTTCTGCTTACCTTACCTG
ACAGCGGCTTACCTTACCTG
AAAAGGTTTACCTTACCTG
GGCCAACTTACCTTACCTG
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TCTGTAAGCTTACCTTACCTG
CTCATTTTACCTTACCTG
TCTGTAAGCTTACCTTACCTG
CTTATTTTACCTTACCTG
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TCTGTAAGCTTACCTTACCTG
TCTGTAAGCTTACCTTACCTG

Ad5/fib16 chimeric fiber

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ATCGCTGGGAGATGGGTTGGTAACAAAGGATGATAAACTATGTTTATCGCTGGGAGATGGGTTAATA
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ACAAATCTACCAATGGAACCTCTTTCCACTAAAAGTTACTGTCACTAAACAGACGTATGTTAGC
TTCTGGAATGGCCTATGCTATGAATTTTTTCATGGTCTCTAAATGCAGAGGAAGCCCCGAAACTACC
GAAGTCACTCTCATTACCTCCCCCTCTTTTTTTCTTATATCAGAGAAGATGACTGAATGCATTAG

bimeric f bar

ATCCCGGTGTTTCTCAGAAATGGGCACAGCGGTAAGCAAAAATAAAGGAATATGGCCTTGCA7
ACTAAGCAATATTAAGAACTATGTTTATTCCTTGGGAGATTGGATGGCCTTGT
TGTTCACCTCTTCCCATACCACAAGCGCCAACTCTGUAUTTAAGTGTGAAGAAT
CGACACCTTAACTTTAAATAATCTTACCTATCTATACCAGTGGCTTAAATGAAG
CTTAAAGTTACTTAAAGTTACTTAAAGTTACTTAAAGTTACTTAAAGTTACTTAAAG
TTAAAGTTACTTAAAGTTACTTAAAGTTACTTAAAGTTACTTAAAGTTACTTAAAG

Figure 4D: Sequence of Ad5/fib28 chimeric fiber

ATGTTGTTGCAGATGAAGCGCGCAAGACCGTCTGAAGATACCTTCAACCCCGTGTATCCATATGGCT
ACGCGCGGAATCAGAATATCCCCCTTCCTCACTCCCCCTTTGTTTCTTCCGATGGATTCCAAAACCTT
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AAGTTGGGAGGCGGACTGACGGTGGAAAAAGAGTCTGGAAACTTAACTGTGAACCCTAAGGCTCCCT
TGCAAGTTGCAAGTGGACAATTGGAATTAGCATATGATTCTCCATTTGATGTTAAAAACAATATGCT
TACTCTTAAAGCAGGTACGGCTTAGCAGTTGTAACGAAAGACAATACTGATTTACAACCACTAATG
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AGAAAATGACAGGCGCACTCTATGGACAACCCAGACACATCTCCAAATTGCAAAATGAGTGAAGTC
AAAGACTCAAAGCTTACTCTTATTCTTACAAAATGCGGAAGTCAAATTCTAGGAAGTGTATCTTTGC
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AACCCGTCAATTCTAAAAGCTATGCCAGAAGTCACATATTTGGAAATGTATATATTGCTGCTAAGCC
ATATAATCCAGTGGTTATTAAAATTAGCTTCAATCAAGAGACACAAAACAATTGTGTCTATTCTATA
TCATTTGACTACACTTGCTCTAAAGAGTATACAGGTATGCAATTCGATGTTACATCTTTCACCTTCT
CCTATATCGCCCAAGAATGAATGCATTAG

chimeric fiber

AAGATACCTTCAACCCCGTGTATCCATATGGCT
CCCCCTTTGTTTCTTCCGATGGATTCCAAAACCTT
CCAATCACCCTAATGGGGATGTCTCACTC
CTGGAAACTTAACTGTGAACCCTAAGGCTCCCT
TGATTCTCCATTTGATGTTAAAAACAATATGCT
GACGAAAGACAATACTGATTTACAACCACTAATG
GGCACTGGCACAAGTGCTCACGGTGGAAACCATAG
TGACAAAAATGGAGATTGGTGGCCTGGGATAA
GACACATCTCCAAATTGCAAAATGAGTGAAGTC
GGGGAAGTCTAGGAAGTGTATCTTTGC
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AATGATTCTACTGTGTCTGGAAAATATGAAAATGCTGTTCCGTTTCATGCCTAACATAACAGCTTATA
AACCCGTCAATTCTAAAAGCTATGCCAGAAGTCACATATTTGGAAATGTATATATTGCTGCTAAGCC
ATATAATCCAGTGGTTATTAAAATTAGCTTCAATCAAGAGACACAAAACAATTGTGTCTATTCTATA
TCATTTGACTACACTTGCTCTAAAGAGTATACAGGTATGCAATTCGATGTTACATCTTTCACCTTCT
CCTATATCGCCCAAGAATGAATGCATTAG

Figure 4E: Sequence Ad5/fib40-L chimeric fiber

ATGTTTGTGTGCAGATGAAGACGCGCGCAAGACCGTCTGAAGATACCTTCAACCCCGTGATCCATATGAAC
ACTACAATCCCCCTTGACATTCCATTTATTACACCCCGCTTTGCTTCCCTCCAACGGCTTGCAAGAAAA
ACCTCCGGGAGTCTCTCAGCCTGAAATACACTGATCCACTTACAACCAAAAACGGGGCTTTAACCTTA
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GCGCCCCCTATCACTAAAACCAACAAAATCGTAGGTTTAAATTACACTAAGCCTCTCGCTCTGCAAA
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TCACAGCCTGTTACTATTAATGCAACAACGAACCTTTCTCTCTTAATAGACGCCCCACTTAATGCTG
ACACGGGCACCTCTTCGCCTTCGAAGTGATGCACCTCTGGACTAGTAGACAAAACACTAAAGGTTTT
GTTTTCTAGCCCCCTCTATCTAGATAATAACTTTCTTACACTAGCCATTGAACGCCCCGCTAGCTCTA
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CGTGCAAAACAATTCTCTCTCCTTAGGGGTTAACCCGCTTTTCTCATCACTGACTCTGGATTAGCT
ATGGACTTAGGAGACGGTCTTGCAATTAGGTGGCTCTAAGTTAATAATCAATCTTGGTCCAGGTTTAC
AAATGTCTAATGGAGCTATTACTTTAGCACTAGATGCAGCGCTGCCTTTGCAATATAAAAAACAACCA
ACTTCAACTCAGAATTGGCTCCGCGTCTGCTTTAATTATGAGCGGAGTAAACACAACATTAACGTC
AATGCCAATACCAGCAAAGGTCTTGCTATTGAAAATAACTCACAGTTGTTAAGCTAGGAAACGGTC
TTCGC'TTTGATAGCTGGGGAAGCATAGCTGTCTCACC'TACTACCCTACCCTTACCACCTATGGAC
CACC GCGGACCCGCTCTCCTAACGCCACTTTTTATGAATCACTAGACGCCAAAGTGTGGCTAGTTTTTA
GTAAAATGCAACGGCATGGTTAACGGGACCATATCCATTAAAGCTCAAAAAGGCACTTTACTTTAAAC
CCACAGCTAGCTTTATTTCTTTTGTCTATGATTTTTTACAGCGACGGAAACGTGGAGGAAAAAC'TATCC
CGTGT'TTGACAACGAAGGGATACTAGCAAACAGTGCCACATGGGGTTATCGACAAGGACAGTCTGCC
AACACTAACGTTTCCAATGCTGTAGAA'TTTATGCC'TAGCTCTAAAAGGTATCCCAATGAAAAAGGTT
CTGAAGTTCAGAACATGGCTCTTACCTACACTTTTTTTGCAAGGTGACCCCTAACATGGCCATATCTTT
TCAGAGCATT'TATAATCATGCAATAGAAGGCTACTCTTAAATTCNCCTGGCGCGTTCGAAATAAT
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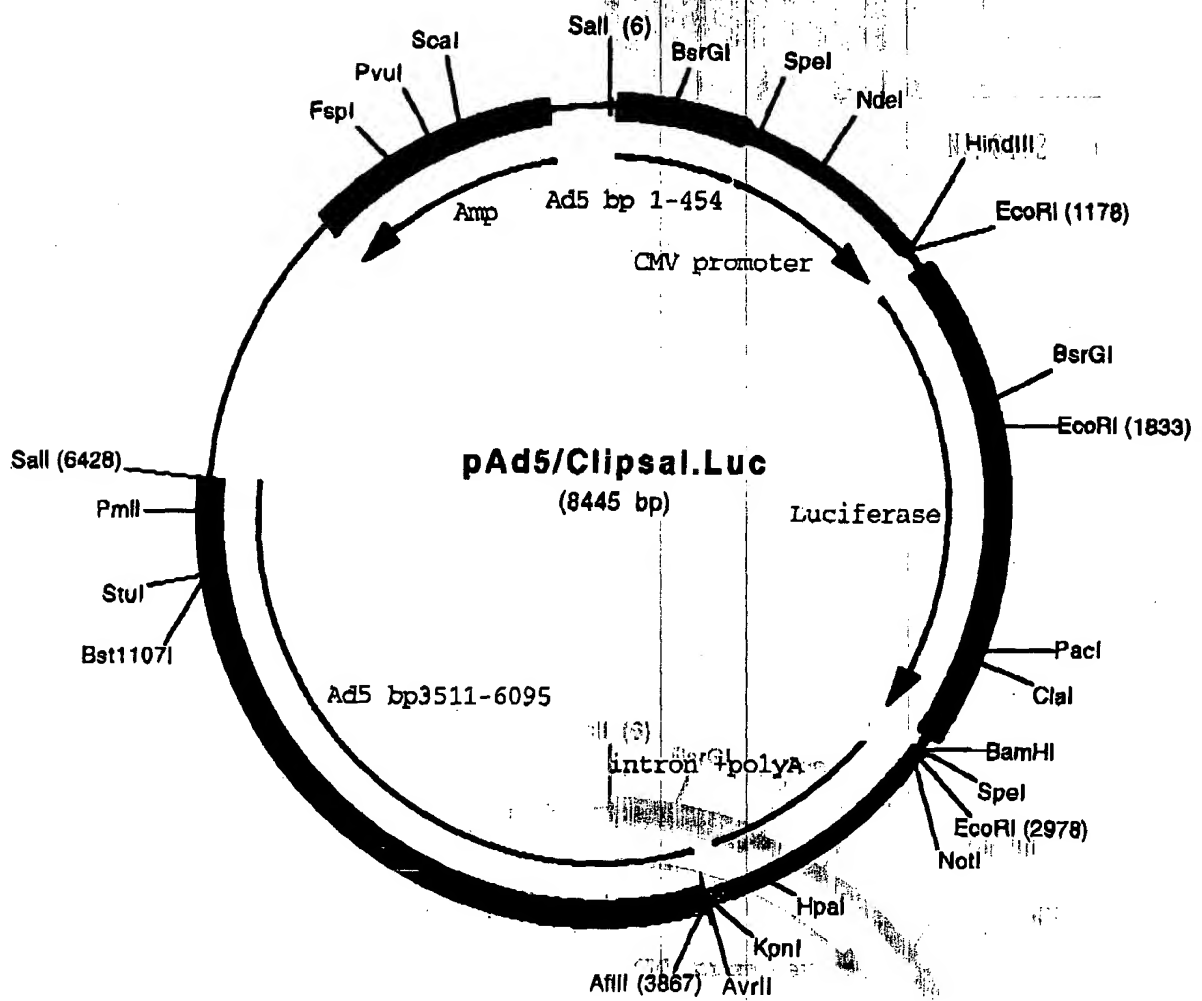


Figure 5

Figure 6: Generation of (chimeric) adenoviruses

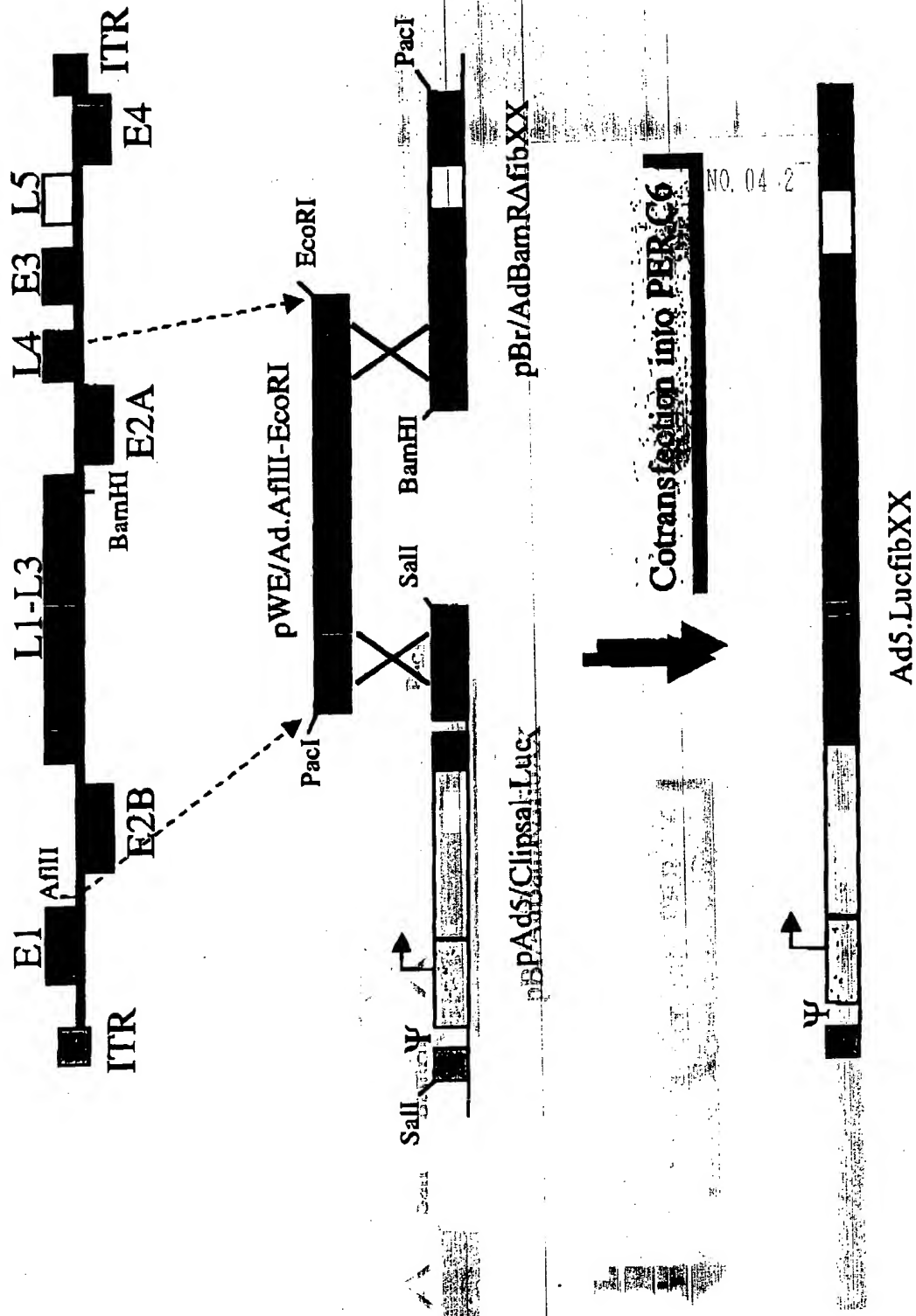


Figure 7a

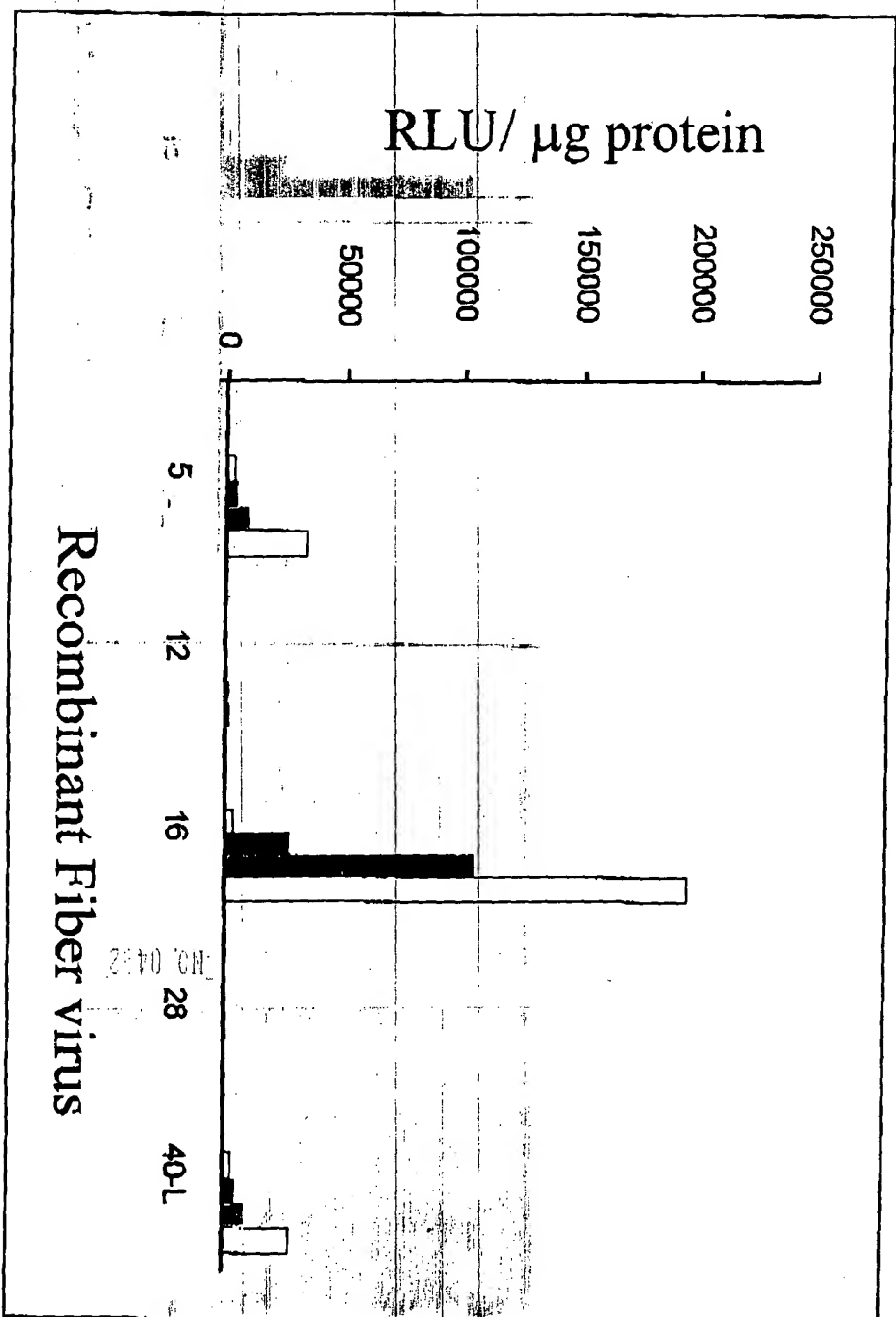


Figure 7b

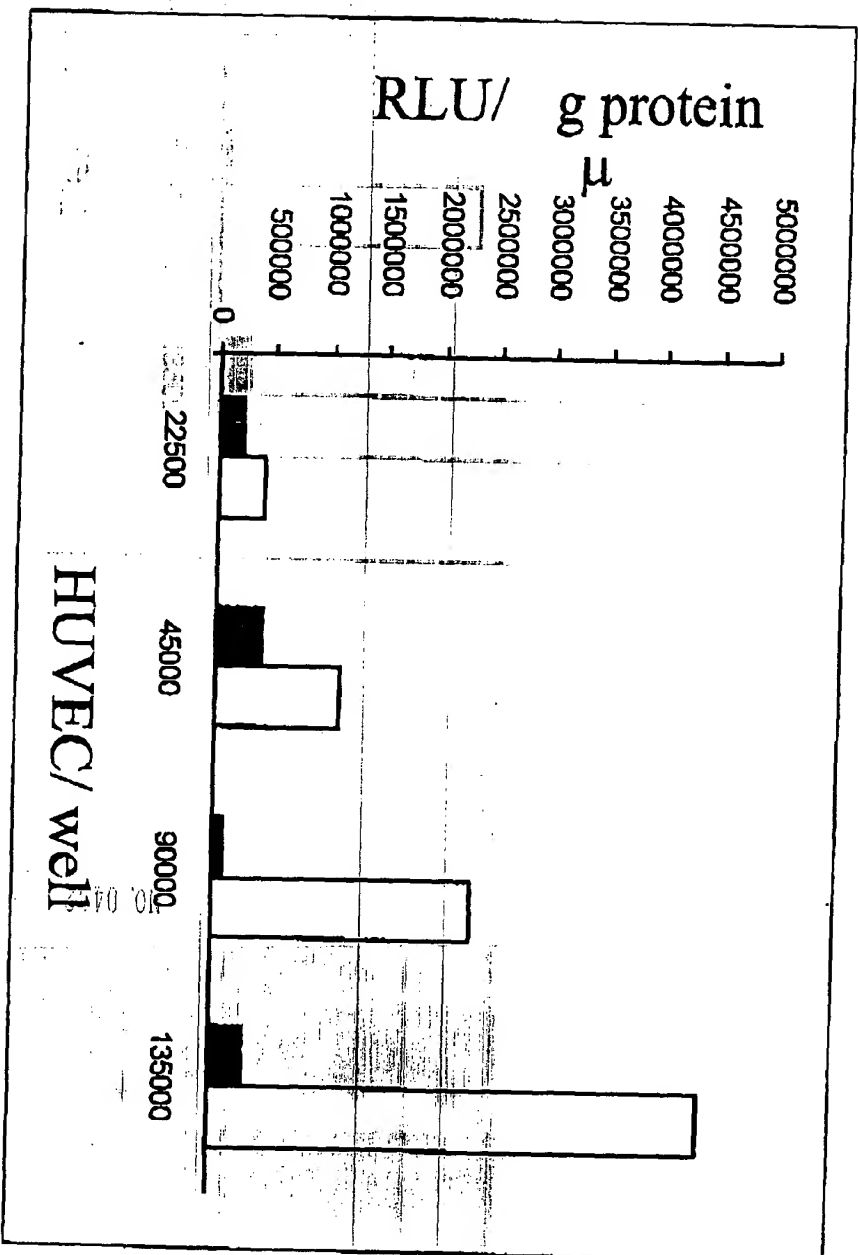


Figure 7c

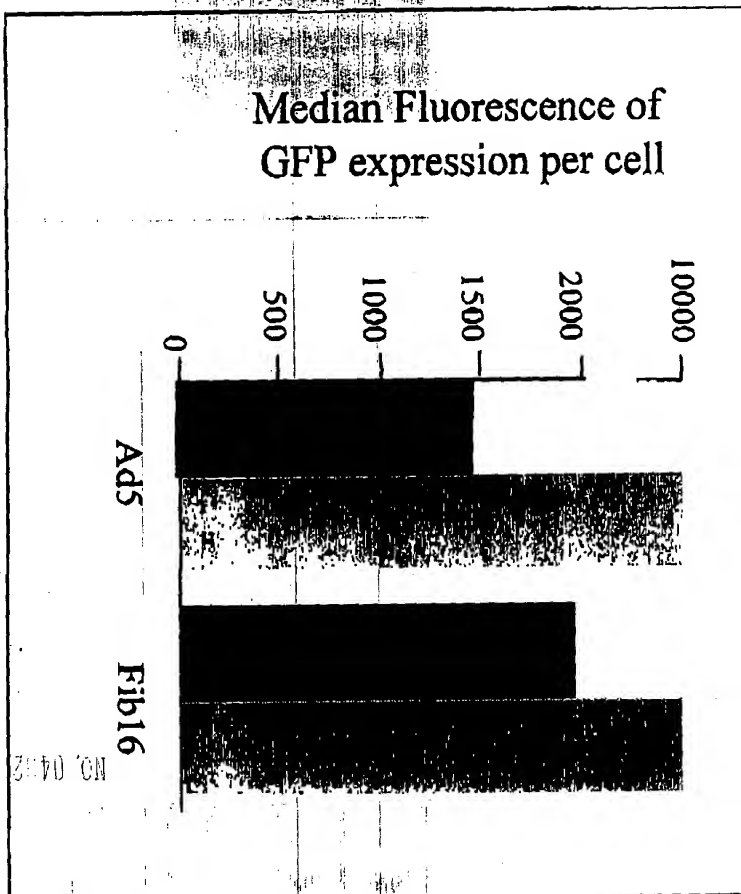


Figure 8a

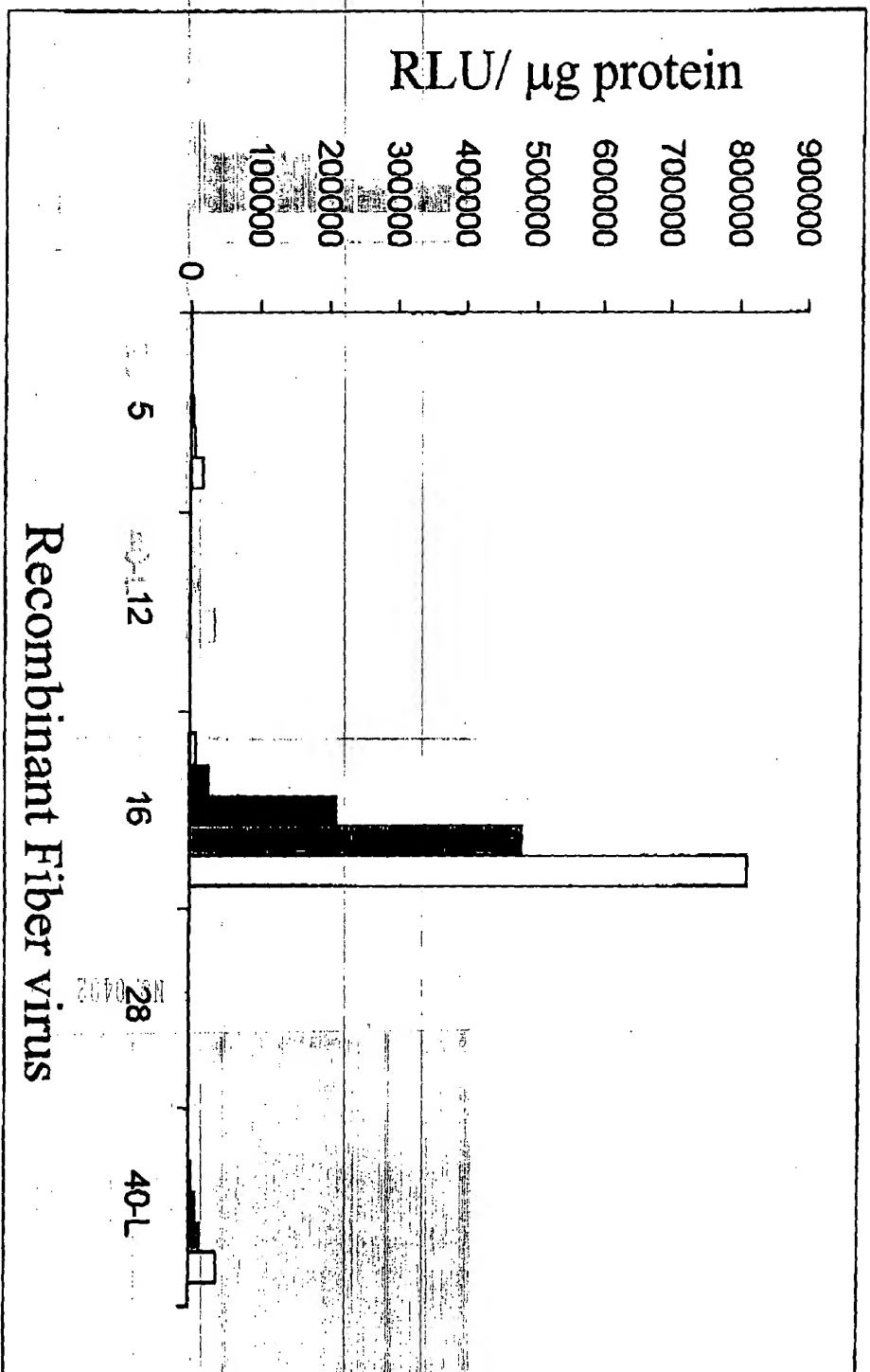
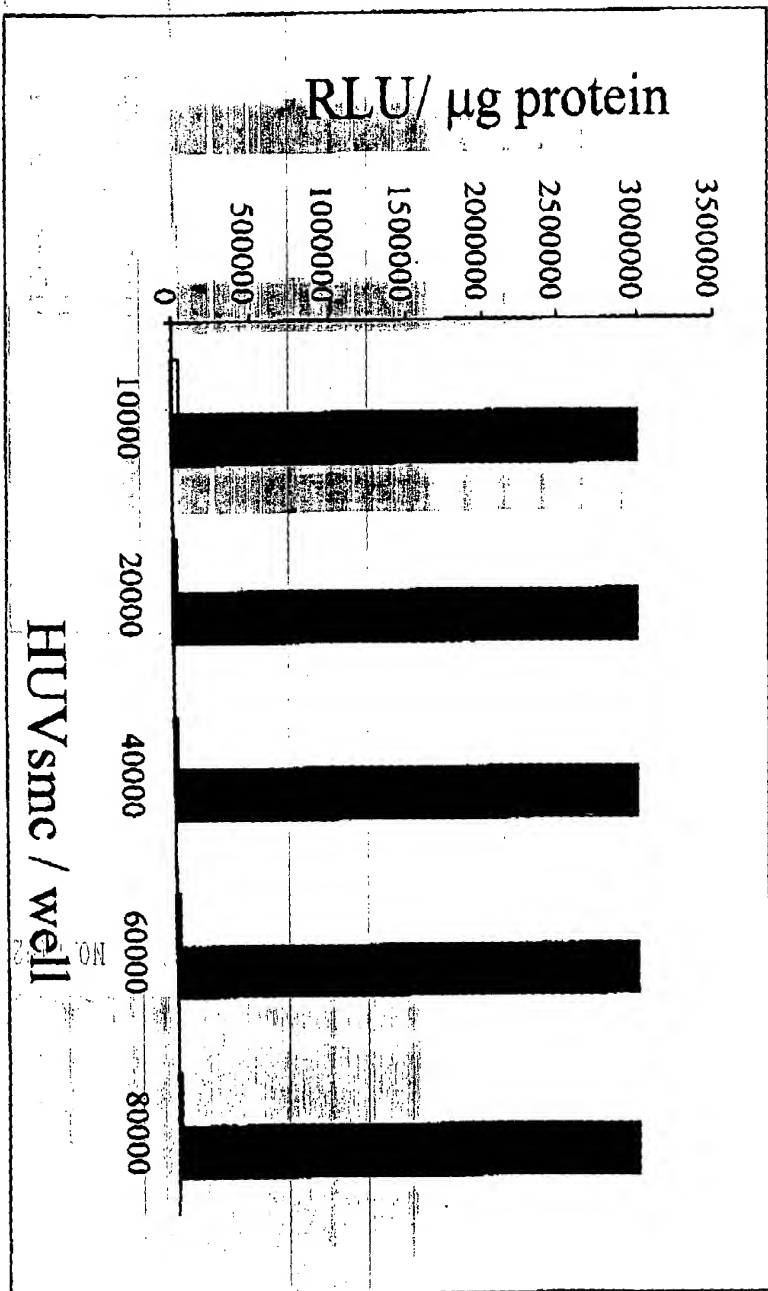
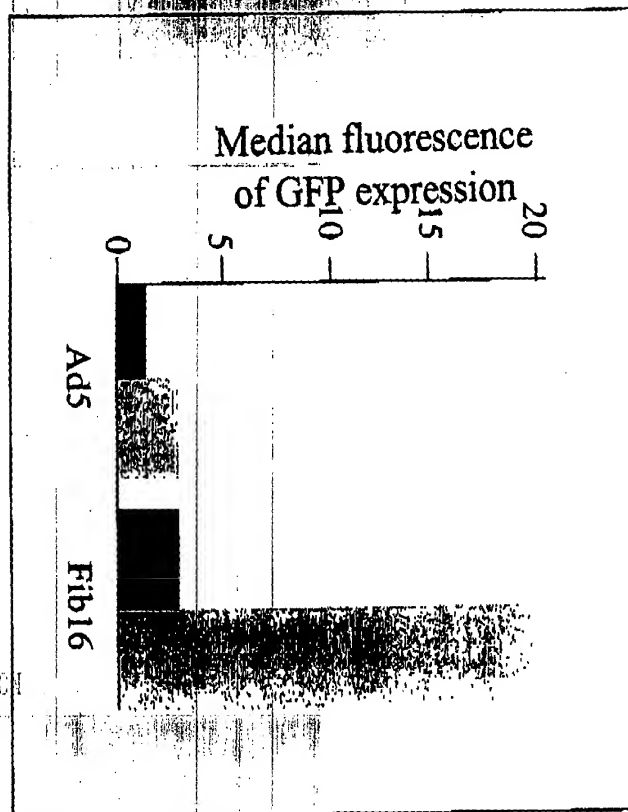
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Figure 8b



10000 20000 40000 60000 80000

Figure 8c



10

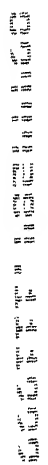


Figure 8e:



Ad5Fiber 16 ntlacZ



Ad5Fiber 16 ntlacZ



Ad5Fiber 5.1 ntlacZ



Negative control

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NO. 04:2

Ad5Fiber 51 ntlacZ

[illegible]

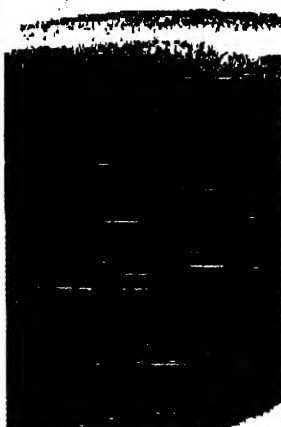
Ad5Fiber 51.ntlacz.

10407

Figure 8g



A Negative control



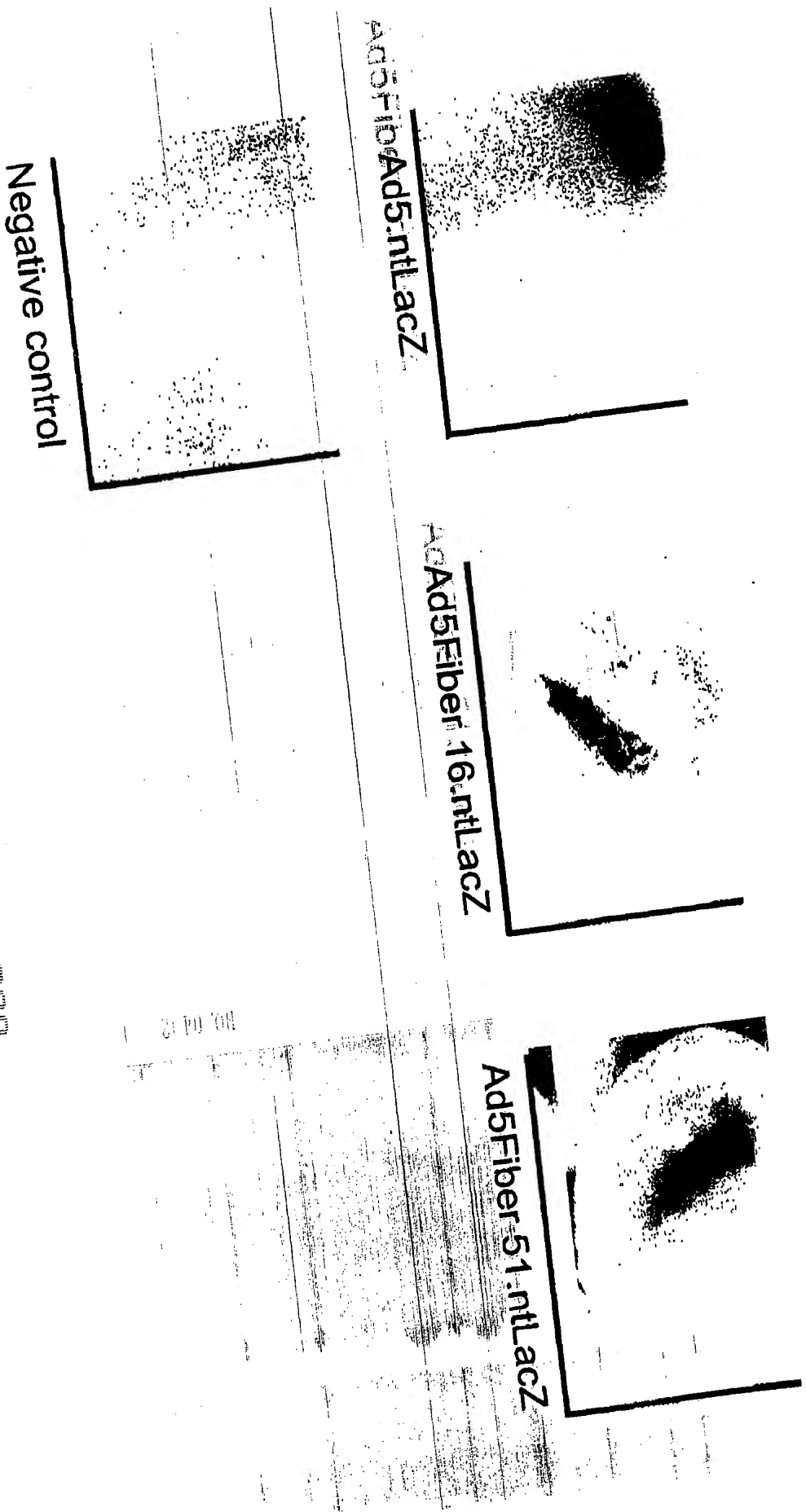
Ad5Fiber 16.nitlacZ



Ad5Fiber 16.nitlacZ

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Figure 8h



[illegible]

Fig 9A.

Alignment Report of Untitled, using Clustal method with Wighted residue weight table.
Thursday, November 19, 1998 18:26

Page 2

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749  AAAACATGGCTACTGGAACCATAACCAGTGCCAAAGGCTT Ad16 genbank.seq
761  AAAACATGGCTACTGGAACCATAACCAGTGCCAAAGGCTT Ad5/fib16.seq

789  CATGCCCAGCACCACCGCCTATCCATTTATAACATACGCC Ad16 genbank.seq
801  CATGCCCAGCACCACCGCCTATCCATTTATAACATACGCC Ad5/fib16.seq

829  ACTGAGACCCTAAATGAAGATTACATTTATGGAGAGTGTT Ad16 genbank.seq
841  ACTGAGACCCTAAATGAAGATTACATTTATGGAGAGTGTT Ad5/fib16.seq

869  ACTACAAATCTACCAATGGAACCTCTCTTTCCACTAAAAGT Ad16 genbank.seq
881  ACTACAAATCTACCAATGGAACCTCTCTTTCCACTAAAAGT Ad5/fib16.seq

909  TACTGTCACTAAACAGACGTATGTTAGCTTCTGGAATG Ad16 genbank.seq
921  TACTGTCACTAAACAGACGTATGTTAGCTTCTGGAATG Ad5/fib16.seq

949  GCCTATGCTATGAATTTTTCATGGTCTCTAAATGCAGAGG Ad16 genbank.seq
961  GCCTATGCTATGAATTTTTCATGGTCTCTAAATGCAGAGG Ad5/fib16.seq

989  AAGCCCCGGAAACTACCGAAGTCACTCTCATTACCTCCCC Ad16 genbank.seq
1001 AAGCCCCGGAAACTACCGAAGTCACTCTCATTACCTCCCC Ad5/fib16.seq

1029 CTTCTTTTTTTTCTTATATCAGAGAGAGATGACTGA Ad16 genbank.seq
1041 CTTCTTTTTTTTCTTATATCAGAGAGAGATGACTGA Ad5/fib16.seq

```

Decoration 'Decoration #1': Box residues that differ from Ad16 genbank.seq

```

CCATTTT...
CCATTTT...

```

```

ACATTTT...
ACATTTT...

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```

TTCTTTT...
TTCTTTT...

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```

TTCTTTT...
TTCTTTT...

```

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TTCTTTT...
TTCTTTT...

```

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TTCTTTT...
TTCTTTT...

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```

TTCTTTT...
TTCTTTT...

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Ad16 genbank.seq

Fig 9B

Alignment Report of Unfiled, using Clustal method with PAM250 residue weight table.
Thursday, November 19, 1998 18:09

Page 1

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1  MAKRRRLSS-SFNPVYPYEDSSSQHPFIN Ad16 fiber protein GenBank
1  M-KRRAR-E-S-E-D-TFNPVYPYEDSSSQHPFIN Ad16A fib protein

30  PGFISSNGFAQSPDGVLTCLKVNPLTTASG Ad16 fiber protein GenBank
30  PGFISSNGFAQSPDGVLTCLKVNPLTTASG Ad16A fib protein

60  PLQLKVGSSSLTVDTIDGSLEENITAAAPLT Ad16 fiber protein GenBank
60  PLQLKVGSSSLTVDTIDGSLEENITAEAPLT Ad16A fib protein

90  KTNHSIGLLIGSGLQTRDDKLCLSLGDGLV Ad16 fiber protein GenBank
90  KTNHSIGLLIGSGLQTRDDKLCLSLGDGLV Ad16A fib protein

120 TKDDKLCLSLGDGLITKNDVLCAKLGHGLV Ad16 fiber protein GenBank
120 TKDDKLCLSLGDGLITKNDVLCAKLGHGLV Ad16A fib protein
      100% ID EV

150 FDSSNAITIENTLWTGAKPSANCVIKEGE Ad16 fiber protein GenBank
150 FDSSNAITIENTLWTGAKPSANCVIKEGE Ad16A fib protein

180 DSPDCKLTLVLVKNGGLINGYITLMGASEY Ad16 fiber protein GenBank
180 DSPDCKLTLVLVKNGGLINGYITLMGASEY Ad16A fib protein

210 TNTLFKNNQVTIDVNLAFDNTGQITLTLSS Ad16 fiber protein GenBank
210 TNTLFKNNQVTIDVNLAFDNTGQIITLTLSS Ad16A fib protein
      100% ID EV

240 LKSNLNFKDNQNMATGTITSAGKFMPS-TTAA Ad16 fiber protein GenBank
240 LKSNLNFKDNQNMATGTITSAGKFMPS-TTAA Ad16A fib protein
      100% ID EV

270 YPFITYATETLNEDYIYGECYKSTNGTLF Ad16 fiber protein GenBank
270 YPFITYATETLNEDYIYGECYKSTNGTLF Ad16A fib protein
      100% ID EV

300 PLKVTVTNLNRRLASGMAYAMNFSW-S-LNAE Ad16 fiber protein GenBank
300 PLKVTVTNLNRRLASGMAYAMNFSW-S-LNAE Ad16A fib protein
      100% ID EV

330 EAPETTEVTLITSPFFFSYIREDD- Ad16 fiber protein GenBank
330 EAPETTEVTLITSPFFFSYIREDD- Ad16A fib protein
      100% ID EV

```

Decoration 'Decoration #1': Box residues that differ from the Consensus.

```

      100% ID EV
      100% ID EV

100% ID EV
100% ID EV

100% ID EV
100% ID EV

100% ID EV
100% ID EV

100% ID EV
100% ID EV

100% ID EV
100% ID EV

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